

**Amendments to the Specification:**

Please substitute the paper copy and computer readable copy of the Sequence Listing presently on file, with the paper copy and computer readable copy of the Sequence Listing submitted herewith.

Please replace paragraph [0137] of the application as published, with the following amended paragraph:

**Fig. 7** is a CLUSTAL W (1.82) multiple sequence alignment for some representative CDK B-type sequences. In bold is a motif found in B-type CDKs; underlined is a catalytic kinase domain and in italics is shown a T-loop activation kinase domain (Magyar *et al.*, 1997 (Plant Cell 9 (2), 223 – 235)). CDKB2.1 is SEQ ID NO:22; CDKB2.2 is SEQ ID NO:23; CDKB1.1 is SEQ ID NO:24; CDKB1.2 is SEQ ID NO:25.

Please replace paragraph [0010] of the application as published, with the following amended paragraph:

[0010] A "B-type CDK nucleic acid" as defined herein is a nucleic acid/gene encoding a protein having: (i) a PPTALRE (SEQ ID NO:6) motif with no mismatches or with a mismatch at position 2 and/or 4 from left to right; (ii) a catalytic kinase domain; and (iii) a T-loop activation kinase domain (Magyar *et al.*, 1997 (Plant Cell 9 (2), 223-235)).